

In the Specification

Please substitute the following paragraphs on page 3, beginning at line 26 through page 4, line 4:

Figure 1: alignment of the SCS0009 ORF (SEQ ID NO:2) with known related polypeptide sequences PREF-1 (SEQ ID NO:38).

Figure 2: Clustal W alignment of predicted amino acid sequences of SCS0009 (SEQ ID NO:2) prediction and splice variants SV3 (SEQ ID NO:39), SV4 (SEQ ID NO:12) and SV5 (SEQ ID NO:40).

Figure 3: Clustal W alignment of predicted amino acid sequences of SCS0009 (SEQ ID NO:2) with SV1 (SEQ ID NO:5) and SV2 (SEQ ID NO:6).

Figure 4: Nucleotide sequence of SCS0009 (SEQ ID NO:1) prediction with translation (SEQ ID NO:2).

Figure 5: Nucleotide sequence with translation of cDNA insert in image clone 5478078 (SCS0009-SV3) (SEQ ID NO:41).

Figure 6 : Alignment of predicted amino acid sequence of SCS0009 (SEQ ID NO:2) with SCS0009-SV3 (SEQ ID NO:8).

Please substitute the following paragraphs on page 4, beginning at line 11:

Figure 13: Nucleotide sequence of SCS0009 (SEQ ID NO:1) prediction with translation (SEQ ID NO:2).

Figure 14: Nucleotide sequence with translation of cDNA insert in image clone 3349698 (SCS0009-SV4) (SEQ ID NO:42).

Figure 15: Alignment of predicted amino acid sequence of SCS0009 (SEQ ID NO:2) with SCS0009-SV4 (SEQ ID NO:12).

Please substitute the following paragraphs on page 4, beginning at line 22:

Figure 22: Nucleotide sequence of SCS0009 (SEQ ID NO:1) prediction with translation (SEQ ID NO:2).

Figure 23: Nucleotide sequence with translation of SCS0009-SV5 (SEQ ID NO:43) PCR product indicating the positions of the SCS0009-AP1, -AP2, -AP3 and -AP4 primers used to generate the SCS0009 sequence.

Figure 24: Nucleotide sequence and translation of cloned SCS0009 ORF (SEQ ID NO:44).

Please substitute the following paragraph on page 28, beginning at line 13:

Van Limpt showed that DLK1 is highly expressed in a subset of neuroblastoma cell lines (Int. J. Cancer. 2003 May 20;105(1):61-9; see also van Limpt V et al. Med Pediatr Oncol. 2000 Dec;35(6):554-8; and Online Mendelian Inheritance in Man<sup>TM</sup> (OMIM)\*176290: [http://www. See Worldwide Website: ncbi.nlm.nih.gov/entrez/query.fcgi?db=OMIM](http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?db=OMIM) ). As such SCS0009 nucleic acid molecules, polypeptides, agonists and antagonists thereof may be useful in diagnosing or treating neuroblastoma, pheochromocytoma, lung tumors, neuroendocrine tumors.

Please substitute the following paragraph on page 38, beginning at line 1:

**Table III      SCS0009-SV3 cloning and sequencing primers**

<b>Primer</b>	<b>Sequence (5'-3')</b>
SCS0009SV3-EX1	AA GCA GGC TTC <u>GCC ACC</u> ATG CCC AGC GGC TGC CGC TG (SEQ ID NO:19)
SCS0009SV3-EX2	GTG ATG GTG ATG GTG CAG TGC TGT GGT CTT TCC AG (SEQ ID NO:20)
GCP Forward	G GGG ACA AGT TTG TAC AAA AAA GCA GGC TTC <u>GCC ACC</u> (SEQ ID NO:21)
GCP Reverse	GGG GAC CAC TTT GTA CAA GAA AGC TGG GTT <b>TCA</b> ATG GTG ATG GTG ATG GTG (SEQ ID NO:22)
SCS0009SV3-SP1	TGA TGC GGC CTT GTG CTA AC (SEQ ID NO:23)
pEAK12F	GCC AGC TTG GCA CTT GAT GT (SEQ ID NO:24)
pEAK12R	GAT GGA GGT GGA CGT GTC AG (SEQ ID NO:25)
21M13	TGT AAA ACG ACG GCC AGT (SEQ ID NO:26)
M13REV	CAG GAA ACA GCT ATG ACC (SEQ ID NO:27)
T7 primer	TAA TAC GAC TCA CTA TAG GG (SEQ ID NO:28)
SP6 primer	ATT TAG GTG ACA CTA TAG (SEQ ID NO:29)

Underlined sequence = Kozak sequence

**Bold** = Stop codon

*Italic* sequence = His tag

Please substitute the following paragraph on page 38, beginning at line 1:

**Table IV      SCS0009-SV4 cloning and sequencing primers**

<b>Primer</b>	<b>Sequence (5'-3')</b>
SCS0009SV4-EX1	AA GCA GGC TTC <u>GCC ACC</u> ATG CCC AGC GGC TGC CGC TG (SEQ ID NO:30)
SCS0009SV4-EX2	<i>GTG ATG GTG ATG GTG</i> GGG TCC AGC CTT GCG CTC GC (SEQ ID NO:31)
GCP Forward	G GGG ACA AGT TTG TAC AAA AAA GCA GGC TTC <u>GCC ACC</u> (SEQ ID NO:21)
GCP Reverse	GGG GAC CAC TTT GTA CAA GAA AGC TGG GTT <b>TCA</b> ATG <i>GTG</i> <i>ATG GTG ATG GTG</i> (SEQ ID NO:22)
pEAK12F	GCC AGC TTG GCA CTT GAT GT (SEQ ID NO:24)
pEAK12R	GAT GGA GGT GGA CGT GTC AG (SEQ ID NO:25)
21M13	TGT AAA ACG ACG GCC AGT (SEQ ID NO:26)
M13REV	CAG GAA ACA GCT ATG ACC (SEQ ID NO:27)
T7 primer	TAA TAC GAC TCA CTA TAG GG (SEQ ID NO:28)
SP6 primer	ATT TAG GTG ACA CTA TAG (SEQ ID NO:29)

Underlined sequence = Kozak sequence

**Bold** = Stop codon

*Italic* sequence = His tag

Please substitute the following paragraph on page 48, beginning at line 5:

**Table V SCS0009 cloning and sequencing primers**

<b>Primer</b>	<b>Sequence (5' -3')</b>
SCS0009-AP1	ACC ATG CCC AGC GGC TGC CGC TGC CTG CAT CTC G <u>(SEQ ID NO:32)</u>
SCS0009-AP2	<u>AGT CAC GCC CAT GGA AGC</u> CTT TGT CAC AGA ACT TGC <u>(SEQ ID NO:33)</u>
SCS0009-AP3	<u>GCA AGT TCT GTG ACA AAG</u> GCT TCC ATG GGC GTG ACT GC <u>(SEQ ID NO:34)</u>
SCS0009-AP4	TCA CAG TGC TGT GGT CTT TCC AGG CTC AGG GGG CAA GTC <u>(SEQ ID NO:35)</u>
SCS0009-EX1	AA GCA GGC TTC <u>GCC ACC</u> ATG CCC AGC GGC TGC CGC TG <u>(SEQ ID NO:36)</u>
SCS0009-EX2	<i>GTG ATG GTG ATG GTG</i> CAG TGC TGT GGT CTT TCC AG <u>(SEQ ID NO:37)</u>
GCP Forward	G GGG ACA AGT TTG TAC AAA AAA GCA GGC TTC <u>GCC ACC</u> <u>(SEQ ID NO:21)</u>
GCP Reverse	GGG GAC CAC TTT GTA CAA GAA AGC TGG GTT <b>TCA</b> ATG <i>GTG ATG GTG ATG GTG</i> <u>(SEQ ID NO:22)</u>
SCS0009-SP1	TGA TGC GGC CTT GTG CTA AC <u>(SEQ ID NO:23)</u>
pEAK12F	GCC AGC TTG GCA CTT GAT GT <u>(SEQ ID NO:24)</u>
pEAK12R	GAT GGA GGT GGA CGT GTC AG <u>(SEQ ID NO:25)</u>
21M13	TGT AAA ACG ACG GCC AGT <u>(SEQ ID NO:26)</u>
M13REV	CAG GAA ACA GCT ATG ACC <u>(SEQ ID NO:27)</u>
T7 primer	TAA TAC GAC TCA CTA TAG GG <u>(SEQ ID NO:28)</u>
SP6 primer	ATT TAG GTG ACA CTA TAG <u>(SEQ ID NO:29)</u>

Double underlined Sequence in red = overlap with adjacent exon.

Underlined sequence = Kozak sequence

**Bold** = Stop codon

*Italic* sequence = His tag

Please replace original Figures 3 and 23 with the attached new Figures 3 and 23.

Please substitute pages 1-28 (Sequence Listing) with the accompanying Sequence Listing (pages 1-37).